



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/050,000 A

Source: PCT

Date Processed by STIC: 11/5/03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebs/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

## Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: <u>10/050,000 A</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <u>    </u> Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <u>    </u> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <u>    </u> Misaligned Amino Numbering	The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <u>    </u> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <u>    </u> Variable Length	Sequence(s) <u>        </u> contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <u>    </u> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) <u>        </u> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <u>    </u> Skipped Sequences (OLD RULES)	Sequence(s) <u>        </u> missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped  Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <u>    </u> Skipped Sequences (NEW RULES)	Sequence(s) <u>        </u> missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <u>  /  </u> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <u>    </u> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <u>    </u> Use of <220>	Sequence(s) <u>        </u> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <u>    </u> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <u>    </u> Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>	



PCT

## RAW SEQUENCE LISTING

DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

3 <110> APPLICANT: NAKARI-SETALA, Tiina  
 5 <120> TITLE OF INVENTION: A METHOD FOR DECREASING THE FOAM FORMATION DURING  
 CULTIVATION OF A  
 6 MICROORGANISM  
 8 <130> FILE REFERENCE: 0365-0529P  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/050,000A  
 C--> 11 <141> CURRENT FILING DATE: 2002-07-10  
 13 <160> NUMBER OF SEQ ID NOS: 8  
 15 <170> SOFTWARE: PatentIn Ver. 2.1  
 17 <210> SEQ ID NO: 1  
 18 <211> LENGTH: 2868  
 19 <212> TYPE: DNA  
 20 <213> ORGANISM: Trichoderma reesei  
 22 <220> FEATURE:  
 23 <221> NAME/KEY: gene  
 24 <222> LOCATION: (1523)..(1950)  
 25 <223> OTHER INFORMATION: hfb1  
 27 <400> SEQUENCE: 1  
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 29 gggacaccga gttgcatatc tcttgatcat tcaaacgaca agtgtgaggt aggcaatcct 120  
 30 cgtatcccat tgctgggctg aaagcttcac acgtatcgca taagcgtctc caaccagtgc 180  
 31 ttagggtgacc cttaaaggata cttacagtaa gactgtatta agtcagtac tctttcactc 240  
 32 gggctttgaa tacgatctc aatactcccg ataacagtaa gaggatgata cagcctgcag 300  
 33 ttggcaaatg taagcgtaat taaactcagc tgaacggccc ttggtgaaag tctctctcga 360  
 34 tcaaagcaaa gctatccaca gacaagggtt aagcaggctc actcttccta cgccttggat 420  
 35 atgcagcttg gccagcatcg cgcattggcca atgatgcacc cttcacggcc caacggatct 480  
 36 cccgttaaac tcccctgtaa cttggcatca ctcatctgtg atcccaacag actgagttgg 540  
 37 gggctgcggc tggcggatgt cggagcaaag gatcacttca agagcccaga tccggttgg 600  
 38 ccattgccaa tggatctaga ttccggacct tgatctcgat cactgagaca tgggtgagttg 660  
 39 cccggacgca ccacaactcc cctgtgtca ttgagtcgcc atatgcgtct tctcagcgtg 720  
 40 caactctgag acggattagt cctcacgatg aaattaactt ccagcttaag ttcgtagcct 780  
 41 tgaatgagtg aagaaatttc aaaaacaaac tgagtagagg tcttgagcag ctggggtgg 840  
 42 acgcccctcc tgcactcttg ggacatcgta cggcagagaa tcaacggatt cacacctttg 900  
 43 ggtcgagatg agctgatctc gacagatacg tgcctcacca cagctgcagc tacctttgcc 960  
 44 caaccattgc gttccaggat cttgatctac atcacgcgag caccgcagcc aggacggaga 1020  
 45 gaacaatccg gccacagagc agcaccgcct tccaactctg ctctggcaa cgtcacacaa 1080  
 46 cctgatatta gatatccacc tgggtgattg ccattgcaga gagggtggcag ttggtgatac 1140  
 47 cgactggcca tgcaagacgc ggccgggcta gctgaaatgt ccccgagagg acaattggga 1200  
 48 gcgtctatga cggcgtggag acgacgggaa aggactcagc cgtcatgttg tgttgccaat 1260  
 49 ttgagattgt tgaccgggaa aggggggacg aagaggatgg ctgggtgagg tggattggg 1320  
 50 aggatgcata attcgactca gtgagcgatg tagagctcca agaataataa tatcccttct 1380  
 51 ctgtcttctc aaaatctcct tccatcttgt ccttcacag caccagagcc agcctgaaca 1440  
 52 cctccagtca acttccctta ccagtacatc tgaatcaaca tccattcttt gaaatctcac 1500  
 53 cacaaccacc atcttcttca aaatgaagtt cttcgccatc gccgtctctt ttgccgccgc 1560

**Does Not Comply  
 Corrected Diskette Needed**

(pg. 3)

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DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

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55 cctcttcagc aacccccagt gctgtgccac ccaagtcctt ggctcatcg gccttgactg 1680
56 caaagtccgt aagttgagcc ataacataag aatcctcttg acggaaatat gccttctcac 1740
57 tcctttaccc ctgaacagcc tcccagaacg ttacgacgg caccgacttc cgcaacgtct 1800
58 gcgcaaaaac cggcgcccag cctctctgct gcgtggcccc cggttgtaagt tgatgcccc 1860
59 gctcaagctc cagtcttttg caaacccatt ctgacacca gactgcaggc cggccagggt 1920
60 cttctgtgcc agaccgccgt cgggtgcttg gatgcccgcc cggggtcaag gtgtgcccgt 1980
61 gagaaagccc acaaagtgtt gatgaggacc atttcggta ctgggaaagt tggctccacg 2040
62 tgtttgggca ggtttgggca agttgtgtag atattccatt cgtacgcat tcttattctc 2100
63 caatatttca gtacactttt cttcataaat caaaaagact gctattctct ttgtgacatg 2160
64 ccggaaggga acaattgctc ttggtctctg ttatttgcaa gtaggagtgg gagattcgcc 2220
65 ttagagaaag tagagaagct gtgcttgacc gtggtgtgac tcgacgagga tggactgaga 2280
66 gtgttaggat taggtcgaa gttgaagtgt atacaggatc gtctggcaac ccacggatcc 2340
67 tatgacttga tgcaatggtg aagatgaatg acagtgtgaa aggaaaagga aatgtccgcc 2400
68 ttcagctgat atccacgcca atgatacagc gatatacctc caatatctgt gggaacgaga 2460
69 catgacatat ttgtgggaac aacttcaaac agcgagccaa gacctcaata tgcacatcca 2520
70 aagccaaaca ttggcaagac gagagacagt cacattgtcg tcgaaagatg gcatcgtaac 2580
71 caaatcatca gctctcatta tcgcctaaac cacagattgt ttgccgtccc ccaactccaa 2640
72 aacgttacta caaaagacat gggcgaatgc aaagacctga aagcaaacc tttttgcgac 2700
73 tcaattccct cctttgtcct cggaatgatg atccttcacc aagtaaaaga aaaagaagat 2760
74 tgagataata catgaaaagc acaacggaaa cgaaagaacc aggaaaagaa taaatctatc 2820
75 acgcaccttg tccccacact aaaagcaaca gggggggtaa aatgaaat 2868
78 <210> SEQ ID NO: 2
79 <211> LENGTH: 3585
80 <212> TYPE: DNA
81 <213> ORGANISM: Trichoderma reesei
83 <220> FEATURE:
84 <221> NAME/KEY: gene
85 <222> LOCATION: (1191)..(1593)
86 <223> OTHER INFORMATION: hfb2
88 <400> SEQUENCE: 2
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90 aactgccgt cgagctctct cttccgacct tggccagctg gacaagcaac acaccaatga 120
91 cgctttgtat tattagagta tatgcaagtc tcaggactat cgactcaact ctaccacccg 180
92 aggacgatcg cggcacgata cgccctcggt ctcatggccc caagcagacc aactgccct 240
93 ggagcaagat tcagcccaag ggagatggac ggcagggcac gccaggcccc caccaccaag 300
94 cactccctt tggccaaatc agcttgcacg tcaagagaca tcgagctgtg cttgaaatt 360
95 actaacaacc agggatggga aacgaagcct gcttttgga agacaacaat gagagagaga 420
96 gagagagggg gagagacaat gagtgccaca aacctggtag tgctccgcca atgcgtctga 480
97 aatgtcacat ccgagtcctt gggcctctgt gagaatgtcc agagtaatac gtgttttgcg 540
98 aatagtcctc tttcttgagg actggatacc tacgataccc tttttgagtt gatgcgggtg 600
99 tttcgaagta ttatctggag gatagaagac gtctaggtaa ctacacaaaa ggcctatact 660
100 ttggggagta gcccaacgaa aggtaactcc tacggcctct tagagccgtc atagatccta 720
101 cagcctcttg gagcgtcat agatcacatc tgtgtagacc gacattctat gaataatcat 780
102 ctcatcatgg ccacatacta ctacatacgt gtctctgcct acctgacatg tagcagtggc 840
103 caagacacca aggccccagc atcaagcctc cctacatc cttccattg tacagcggca 900
104 gagagattgc gatgagcct ctccctacct acagcgggt gacaatgtcc gtataccacc 960
105 agccaacgtg atgaaaacaa ggacatgagg aacagcctgc gagagctgga agatgaagag 1020
106 ggccagaaaa aaaagtataa agaagacctc gattcccgcc atccaacaat cttttccatc 1080

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PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

W--&gt;

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107 ctcatcagca cactcatcta caaccatcac cacattcact caactcctct ttctcaactc 1140
108 tccaaacaca aacattcttt gttgaatacc aaccatcacc acctttcaag atgcagttct 1200
109 tcgccgtcgc cctcttcgcc accagcgccc tggctgctgt ctgccctacc ggcctcttct 1260
110 ccaaccctct gtgctgtgcc accaacgtcc tcgacctcat tggcggtgac tgcaagaccc 1320
111 gtatgttgaa ttccaatctc tgggcatcct gacattggac gatacagttg acttacacga 1380
112 tgctttacag ctaccatcgc cgtcgacact ggcgccatct tccaggctca ctgtgccagc 1440
113 aagggtcca agcctctttg ctgcgttget cccgtggtaa gtagtgctcg caatggcaaa 1500
114 gaagtaaaaa gacatttggg cctgggatcg ctaactcttg atatcaaggc cgaccaggct 1560
115 ctctgtgcc agaaggccat cggcaccttc taaagcaatg gcttgcttta ctgccggcag 1620
116 tctttgagaa ctctgggctc aaaaagacg acttgcattg atcatggggg ctgcgaaatg 1680
117 ggaggatttg gaggggattg aggctgggtt tggcctatta gaggattgca taatggaaga 1740
118 tttgcgagca ggacatagac gtatctagag ttctagtcaa tacattattg aaaagttgga 1800
119 gtatacctat cgctggcact ggtatcttga agatatcttc tcttcttgtg aggttatgta 1860
120 tggaatcag tcgaaatcta tttgaagaca gagctcaagg ttcaaacatt cacctgga 1920
121 ttgaccattt tgtttcgatg gttgcagttg gtgggtgtca cttctgcaat catgtacgag 1980
122 cacaagtata gcagtattcc atctgatctg catctgggta aatgtcgcca ctctacctag 2040
123 gtacccaata aataccgaat tggtcagctc tcgggtgaca aaccggcccg cttttcgacc 2100
124 gtgctctgtc caattctagg ctgttcaatg gttcctgact gtgataaacc ttggagctat 2160
125 cataacttac cttacaataa atccaactgc cggcacttgc ttcccttcac ccaaccactc 2220
126 gcaaacatca cgcaacctgt ctcgatcccc tgtccgaaat ctgcttgga acgtatcatc 2280
127 acaaatcata cacacagaca aaaaggagcc aaagcagcaa tggcaagaca ccgaggccgg 2340
128 cagcgcgccc gtcgccgttt ttaaaaagcg aagcgcaaa ggcaaagcca acctgcgcaa 2400
129 acgaacaacg aagccttccc cccgccgcga gcgacagcga cagcgacagc gacttttct 2460
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132 ttcacggcca acagaagcgt cccgattgct gacagcaacg acgcgacca gacagcaac 2640
133 tggtagcagc aggacgcaaa ggacgcgctc tcggcaaaga acctcctcgg atcttcgaga 2700
134 gcgtccaagg acgcgcagcc agacggcacg tacaagggcc tggcgaacca gacgtcctt 2760
135 atacaaaaga atccggatgc gccccggaag acagttgggc ccgtcaaggc gcctaccaac 2820
136 atccgcaccg tcaccattac agattatgcc ccggacacgt gtaaagagtg agtttgcatc 2880
137 aatagccaga atcccccccc ccgataccgt acattgagca tatgctgact cgtcataatc 2940
138 tttctagtta tcgcataacc ggctatataa gtactccct tttccatgat tattccagtc 3000
139 gcgtactgac atttctagga gcctttactg tggttttggc gacaattgca agtatcttca 3060
140 cgcgagagaa gacctcaagg caggctggca gctggatcaa gagggggaaa aggtcaccaa 3120
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142 ggacgagggc gacgacgacg acgacgaaga ggcgatgctc gagaacattc cgtttgcctg 3240
143 catcatctgc agggaatcgt acaaggagcc gattgtgacg aggtgcgggc actacttttg 3300
144 cctgccgtgc gctctgcagc ggtacaagaa ggatccgacg tgtgcggcgt gtggctcggg 3360
145 cacgaatggc gtgtttaatt cggcgacgag gttgaagaag ctgctggaga agaagaggga 3420
146 gagggcggcc aggaggagac aggaggcgat agagaggggc gaggaagtca gtgatgaaga 3480
147 ggaggaggag gaggaggact gatgatgatg gggcagatg acgatgcagg tcgactctag 3540
148 agatccccgg taccgagctc gaattcatcg atgatatcag atccc 3585
151 <210> SEQ ID NO: 3
152 <211> LENGTH: 63
153 <212> TYPE: DNA
154 <213> ORGANISM: Artificial Sequence
156 <220> FEATURE:
157 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer
159 <400> SEQUENCE: 3

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- Please explain "N" locations. See item #9 on error summary sheet and explanation on page 5.

## RAW SEQUENCE LISTING

DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

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160 actacacgga ggagctcgac gacttcgagc agcccgagct gcacgcagag caacggcaac 60
161 ggc 63
164 <210> SEQ ID NO: 4
165 <211> LENGTH: 30
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Description of Artificial Sequence: 3'primer
172 <400> SEQUENCE: 4
173 tcgtacggat cctcaagcac cgacggcgg 30
176 <210> SEQ ID NO: 5
177 <211> LENGTH: 13
178 <212> TYPE: PRT
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence: linker
184 <400> SEQUENCE: 5
185 Pro Gly Ala Ser Thr Ser Thr Gly Met Gly Pro Gly Gly
186 1 5 10
189 <210> SEQ ID NO: 6
190 <211> LENGTH: 14
191 <212> TYPE: PRT
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: linker
197 <400> SEQUENCE: 6
198 Gly Thr Leu Val Pro Arg Gly Pro Ala Glu Val Asn Leu Val
199 1 5 10
202 <210> SEQ ID NO: 7
203 <211> LENGTH: 48
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Description of Artificial Sequence: 5'primer
210 <400> SEQUENCE: 7
211 gaattcggta cctcgtccc tcgcggtccc gccgaagtga acctggtg 48
214 <210> SEQ ID NO: 8
215 <211> LENGTH: 34
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: Description of Artificial Sequence: 3'primer
222 <400> SEQUENCE: 8
223 tgaattccat atgctaaccc cgtttcatct ccag 34

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/050,000A

DATE: 01/05/2004  
TIME: 11:45:57

Input Set : A:\2002-07-07 0365-0529P.ST25.txt  
Output Set: N:\CRF4\01052004\J050000A.raw

Please Note: *Error Explanation:*

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1917,2160,3515

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/050,000A

DATE: 01/05/2004

TIME: 11:45:57

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1860  
M:341 Repeated in SeqNo=2